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## Sugar utilization in the hyperthermophilic, sulfate-reducing archaeon *Archaeoglobus fulgidus* strain 7324: starch degradation to acetate and CO<sub>2</sub> via a modified Embden-Meyerhof pathway and acetyl-CoA synthetase (ADP-forming)

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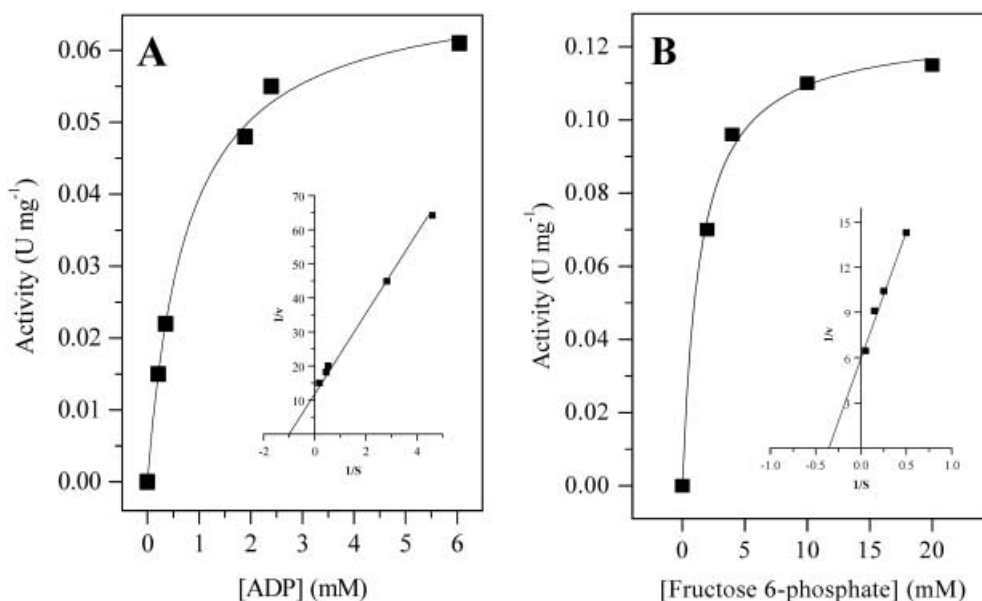
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The Michaelis-Menten plots and the corresponding Lineweaver-Burk plots shown in Fig. 2A and B were unfortunately incorrect. Furthermore, in Fig. 3, the stoichiometric

factors for ADP and ATP in reaction 9, and for ADP, Pi, and ATP in reaction 11, should read 2 instead of 1. The corrected figures appear below:

**Fig. 2A, B** ADP-dependent 6-phosphofructokinase activity at 50°C in cell extracts of starch-grown *A. fulgidus* strain 7324. Rate dependence on **A** the ADP concentration and **B** the fructose 6-phosphate concentration. *Inserts* Double reciprocal plots of the rates vs the corresponding substrate concentrations



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**Fig. 3** Proposed pathway of starch degradation – via glucose – to acetate and  $\text{CO}_2$  in *A. fulgidus* strain 7324. The enzymes of the modified Embden-Meyerhof pathway (enzymes 1–9) and of pyruvate conversion to acetate and  $\text{CO}_2$  (enzymes 10–11) are shown. 1 ADP-dependent hexokinase, 2 glucose-6-phosphate isomerase, 3 ADP-dependent 6-phosphofructokinase, 4 fructose-1,6-bisphosphate aldolase, 5 triosephosphate isomerase, 6 glyceraldehyde-3-phosphate: ferredoxin oxidoreductase, 7 mutase, 8 enolase, 9 pyruvate kinase, 10 pyruvate:ferredoxin oxidoreductase, 11 ADP-forming acetyl-CoA synthetase.  $Fd_{ox}$  oxidized ferredoxin,  $Fd_{red}$  reduced ferredoxin

